

SEQUENCE LISTING

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<120> METHODS AND COMPOSITIONS FOR MODULATING FERTILITY

<130> 9426-015-999

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<151> 2000-02-01

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<212> DNA

<213> Homo sapiens

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<212> PRT

<213> mouse

<400> 2

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 Val Asp Leu Tyr Ser Pro Lys Gly Gln Gln Asp Gln Asp Arg Lys Val
 20 25 30
 Ile Cys Phe Val Asp Val Ser Thr Leu Asn Val Glu Asp Phe Asn Ser
 35 40 45
 Lys Gly Ala Ala Gly Pro Arg Ser Glu Gly Glu Leu Asn Leu Glu Asn
 50 55 60
 Leu Glu Glu Lys Glu Ile Ile Val Ile Lys Asp Thr Glu Lys Gln Asp
 65 70 75 80
 Gln Ser Lys Thr Glu Gly Ser Val Cys Leu Phe Lys Gln Ala Pro Ser
 85 90 95
 Asp Pro Ile Ser Val Leu Asn Trp Leu Leu Asn Asp Leu Gln Lys Tyr
 100 105 110
 Ala Ser Gly Phe Gln Gln Ala Leu Ser Pro Ser Ala Ser Ser Cys Lys
 115 120 125
 His Lys Val Gly Asp Leu Glu Gly Asp Tyr His Lys Ile Pro Ser Glu
 130 135 140

Glu Cys Tyr Ser Val Tyr Ala Asp Gln Val Asn Leu Asp Tyr Leu Asn
 145 150 155 160
 Lys Gly Pro Gln Asn Leu Arg Leu Glu Met Ala Ala Ser Lys Asn Thr
 165 170 175
 Asn Asn Asn Gln Ser Pro Ser Asn Pro Ala Thr Lys Ser Pro Ser Asn
 180 185 190
 Gln Arg Ser Val Ala Thr Pro Asp Gly Glu Cys Ser Met Asp Asp Leu
 195 200 205
 Ser Tyr Tyr Val Asn Arg Leu Ser Ser Leu Val Ile Gln Met Ala Arg
 210 215 220
 Lys Glu Ile Lys Asp Lys Leu Glu Gly Gly Asn Lys Cys Leu His His
 225 230 235 240
 Ser Met Tyr Thr Ser Gly Glu Lys Gly Lys Thr Ser Pro Arg Ser Ala
 245 250 255
 Val Ser Lys Ile Ala Ser Glu Met Ala His Glu Ala Val Glu Leu Thr
 260 265 270
 Ser Ser Glu Met Arg Gly Asn Gly Glu Glu Gly Arg Asp Gly Arg Lys
 275 280 285
 Thr Phe Leu Tyr Ser Glu Leu Ser Asn Lys Asn Lys Cys Gly Glu Lys
 290 295 300
 Gln Gln Met Cys Pro Lys Asp Ser Lys Glu Phe Ala Asp Ser Ile Ser
 305 310 315 320
 Lys Gly Leu Met Val Tyr Ala Asn Gln Val Ala Ser Asp Met Met Val
 325 330 335
 Ser Val Met Lys Thr Leu Lys Val His Ser Cys Gly Lys Pro Ile Pro
 340 345 350
 Ala Cys Val Val Leu Lys Arg Val Leu Leu Lys His Thr Lys Glu Ile
 355 360 365
 Val Ser Asp Leu Ile Asp Ser Cys Met Lys Asn Leu His Asn Ile Thr
 370 375 380
 Gly Val Leu Met Thr Asp Ser Asp Phe Val Ser Ala Val Lys Arg Asn
 385 390 395 400
 Leu Phe Asn His Gly Lys Gln Asn Ala Ala Asp Ile Met Glu Ala Met
 405 410 415
 Leu Lys Arg Leu Val Ser Ala Leu Leu Gly Glu Lys Lys Glu Thr Lys
 420 425 430
 Ser Gln Ser Leu Ala Tyr Ala Thr Leu Lys Ala Gly Thr His Asp Pro
 435 440 445
 Lys Cys Lys Asn Gln Ser Leu Glu Phe Ser Ala Met Lys Ala Glu Met
 450 455 460
 Lys Gly Lys Asp Lys Gly Lys Thr Lys Gly Asp Pro Cys Cys Lys Ser
 465 470 475 480

Leu Thr Ser Ala Glu Arg Val Ser Glu His Ile Leu Lys Glu Ser Leu
 485 490 495
 Thr Met Trp Asn Asn Gln Lys Gln Gly Thr Gln Gly Arg Val Pro Asn
 500 505 510
 Lys Val Cys Pro Ser Lys Asp Glu Lys Arg Glu Lys Ile Ser Pro Ser
 515 520 525
 Thr Asp Ser Leu Ala Lys Asp Leu Ile Val Ser Ala Leu Met Leu Ile
 530 535 540
 Gln Tyr His Leu Thr Gln Gln Ala Lys Gly Lys Asp Pro Cys Glu Glu
 545 550 555 560
 Glu Cys Pro Gly Ser Ser Met Gly Tyr Met Ser Gln Ser Ala Gln Tyr
 565 570 575
 Glu Lys Ser Gly Gly Gln Ser Ser Lys Ser Leu Ser Met Lys His
 580 585 590
 Phe Glu Ser Arg Gly Ala Pro Gly Pro Ser Thr Cys Ala Lys Glu Asn
 595 600 605
 Gln Leu Glu Ser Gln Lys Met Asp Met Ser Asn Met Val Leu Ser Leu
 610 615 620
 Ile Gln Lys Leu Leu Ser Glu Ser Pro Phe Ser Cys Asp Glu Leu Ser
 625 630 635 640
 Glu Ser Glu Asn Lys Arg Cys Cys Asp Ser Arg Ser Lys Gln Ala Ala
 645 650 655
 Pro Val Ala Lys Arg Pro Glu Asp Gln Ser Gln Asp Ser Thr Glu Met
 660 665 670
 Asp Phe Ile Ser Gly Met Lys Gln Met Asn Arg Gln Phe Ile Asp Gln
 675 680 685
 Leu Val Glu Ser Val Met Lys Leu Cys Leu Ile Met Ala Lys Tyr Ser
 690 695 700
 Asn Asn Gly Ala Ala Leu Ala Glu Leu Glu Glu Gln Ala Ala Leu Ala
 705 710 715 720
 Ser Asn Gly Pro Arg Cys Gly Arg Glu Ala Val Met Ser Gln Ser Tyr
 725 730 735
 Pro Glu Thr Pro Gly Pro Glu Val Ile Val Asn Asn Gln Cys Ser Thr
 740 745 750
 Ser Asn Leu Gln Lys Gln Leu Gln Ala Val Leu Gln Trp Ile Ala Ala
 755 760 765
 Ser Gln Phe Asn Val Pro Met Leu Tyr Phe Met Gly Asp Asp Asp Gly
 770 775 780
 Gln Leu Glu Lys Leu Pro Glu Val Ser Ala Lys Ala Ala Glu Lys Gly
 785 790 795 800
 Tyr Ser Val Gly Asp Leu Leu Gln Glu Val Met Lys Phe Ala Lys Glu
 805 810 815

Thr Gln Leu Asp Glu Ala Val Gly Asn Met Ala Arg Lys Gln Leu Leu
820 825 830

Asp Trp Leu Leu Ala Asn Leu
835

<210> 3

<211> 853

<212> PRT

<213> Homo sapiens

<400> 3

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Val Asp Val Tyr Ser Pro Gly Asp Asn Gln Ala Gln Asp Trp Lys Met
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Asp Thr Ser Thr Asp Pro Val Arg Val Leu Ser Trp Leu Arg Arg Asp
35 40 45

Leu Glu Lys Ser Thr Ala Glu Phe Gln Asp Val Arg Phe Lys Pro Gly
50 55 60

Glu Ser Phe Gly Gly Glu Thr Ser Asn Ser Gly Asp Pro His Lys Gly
65 70 75 80

Phe Ser Val Asp Tyr Tyr Asn Thr Thr Thr Lys Gly Thr Pro Glu Arg
85 90 95

Leu His Phe Glu Met Thr His Lys Glu Ile Pro Cys Gln Gly Pro Arg
100 105 110

Ala Gln Leu Gly Asn Gly Ser Ser Val Asp Glu Val Ser Phe Tyr Ala
115 120 125

Asn Arg Leu Thr Asn Leu Val Ile Ala Met Ala Arg Lys Glu Ile Asn
130 135 140

Glu Lys Ile Asp Gly Ser Glu Asn Lys Cys Val Tyr Gln Ser Leu Tyr
145 150 155 160

Met Gly Asn Glu Pro Thr Pro Thr Lys Ser Leu Ser Lys Ile Ala Ser
165 170 175

Glu Leu Val Asn Glu Thr Val Ser Ala Cys Ser Arg Asn Ala Ala Pro
180 185 190

Asp Lys Ala Pro Gly Ser Gly Asp Arg Val Ser Gly Ser Ser Gln Ser
195 200 205

Pro Pro Asn Leu Lys Tyr Lys Ser Thr Leu Lys Ile Lys Glu Ser Thr
210 215 220

Lys Glu Arg Gln Gly Pro Asp Asp Lys Pro Pro Ser Lys Lys Ser Phe
225 230 235 240

Phe Tyr Lys Glu Val Phe Glu Ser Arg Asn Gly Asp Tyr Ala Arg Glu
 245 250 255
 Gly Gly Arg Phe Phe Pro Arg Glu Arg Lys Arg Phe Arg Gly Gln Glu
 260 265 270
 Arg Pro Asp Asp Phe Thr Ala Ser Val Gly Glu Gly Ile Met Thr Tyr
 275 280 285
 Ala Asn Ser Val Val Ser Asp Met Met Val Ser Ile Met Lys Thr Leu
 290 295 300
 Lys Ile Gln Val Lys Asp Thr Thr Ile Ala Thr Ile Leu Leu Lys Lys
 305 310 315 320
 Val Leu Leu Lys His Ala Lys Glu Val Val Ser Asp Leu Ile Asp Ser
 325 330 335
 Phe Leu Arg Asn Leu His Ser Val Thr Gly Thr Leu Met Thr Asp Thr
 340 345 350
 Gln Phe Val Ser Ala Val Lys Arg Thr Val Phe Ser His Gly Ser Gln
 355 360 365
 Lys Ala Thr Asp Ile Met Asp Ala Met Leu Arg Lys Leu Tyr Asn Val
 370 375 380
 Met Phe Ala Lys Lys Val Pro Glu His Val Arg Lys Ala Gln Asp Lys
 385 390 395 400
 Ala Val Ser Tyr Ser Leu Ile Ser Met Lys Gly Met Gly Asp Pro Lys
 405 410 415
 Asn Arg Asn Val Asn Phe Ala Met Lys Ser Glu Thr Lys Leu Arg Glu
 420 425 430
 Lys Met Tyr Ser Glu Pro Lys Ser Glu Glu Glu Thr Cys Ala Lys Thr
 435 440 445
 Leu Gly Glu His Ile Ile Lys Glu Gly Leu Thr Leu Trp His Lys Ser
 450 455 460
 Gln Gln Asn Glu Cys Lys Ser Leu Gly Phe Gln His Ala Ala Phe Glu
 465 470 475 480
 Ala Pro Asn Thr Gln Arg Lys Pro Ala Ser Asp Ile Ser Phe Glu Tyr
 485 490 495
 Pro Glu Asp Thr Gly Asn Leu Ser Leu Pro Pro Tyr Pro Glu Lys
 500 505 510
 Pro Glu Asn Phe Met Tyr Asp Ser Asp Ser Trp Ala Lys Asp Leu Ile
 515 520 525
 Val Ser Ala Leu Leu Leu Ile Gln Tyr His Leu Ala Gln Gly Gly Arg
 530 535 540
 Arg Asp Ala Arg Ser Phe Val Glu Ala Ala Gly Thr Thr Asn Phe Pro
 545 550 555 560
 Ala Asn Glu Pro Pro Val Ala Pro Asp Glu Ser Cys Leu Lys Ser Ala
 565 570 575

Pro Ile Val Gly Asp Gln Glu Gln Ala Glu Lys Lys Asp Leu Arg Ser
 580 585 590
 Val Phe Phe Asn Ser Ile Arg Asn Leu Leu Ser Glu Thr Ile Phe Lys
 595 600 605
 Arg Asp Gln Ser Pro Glu Pro Lys Val Pro Glu Gln Pro Val Lys Glu
 610 615 620
 Asp Arg Lys Leu Cys Glu Arg Pro Leu Ala Ser Ser Pro Pro Arg Leu
 625 630 635
 Tyr Glu Asp Asp Glu Thr Pro Gly Ala Leu Ser Gly Leu Thr Lys Met
 645 650 655
 Ala Val Ser Gln Ile Asp Gly His Met Ser Gly Gln Met Val Glu His
 660 665 670
 Leu Met Asn Ser Val Met Lys Leu Cys Val Ile Ile Ala Lys Ser Cys
 675 680 685
 Asp Ala Ser Leu Ala Glu Leu Gly Asp Asp Lys Leu Gly Asp Ala Ser
 690 695 700
 Arg Leu Thr Ser Ala Phe Pro Asp Ser Leu Tyr Glu Cys Leu Pro Ala
 705 710 715 720
 Lys Gly Thr Gly Ser Ala Glu Ala Val Leu Gln Asn Ala Tyr Gln Ala
 725 730 735
 Ile His Asn Glu Met Arg Gly Thr Ser Gly Gln Pro Pro Glu Gly Cys
 740 745 750
 Ala Ala Pro Thr Val Ile Val Ser Asn His Asn Leu Thr Asp Thr Val
 755 760 765
 Gln Asn Lys Gln Leu Gln Ala Val Leu Gln Trp Val Ala Ala Ser Glu
 770 775 780
 Leu Asn Val Pro Ile Leu Tyr Phe Ala Gly Asp Asp Glu Gly Ile Gln
 785 790 795 800
 Glu Lys Leu Leu Gln Leu Ser Ala Ala Ala Val Asp Lys Gly Cys Ser
 805 810 815
 Val Gly Glu Val Leu Gln Ser Val Leu Arg Tyr Glu Lys Glu Arg Gln
 820 825 830
 Leu Asn Glu Ala Val Gly Asn Val Thr Pro Leu Gln Leu Leu Asp Trp
 835 840 845
 Leu Met Val Asn Leu
 850

<210> 4

<211> 845

<212> PRT

<213> Homo sapiens

<400> 4

Met Ser Asp Asp Ile Asp Trp Leu Arg Ser His Arg Gly Val Cys Lys
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Val Asp Leu Tyr Asn Pro Glu Gly Gln Gln Asp Gln Asp Arg Lys Val
20 25 30

Ile Cys Phe Val Asp Val Ser Thr Leu Asn Val Glu Asp Lys Asp Tyr
35 40 45

Lys Asp Ala Ala Ser Ser Ser Ser Glu Gly Asn Leu Asn Leu Gly Ser
50 55 60

Leu Glu Glu Lys Glu Ile Ile Val Ile Lys Asp Thr Glu Lys Lys Asp
65 70 75 80

Gln Ser Lys Thr Glu Gly Ser Val Cys Leu Phe Lys Gln Ala Pro Ser
85 90 95

Asp Pro Val Ser Val Leu Asn Trp Leu Leu Ser Asp Leu Gln Lys Tyr
100 105 110

Ala Trp Gly Phe Gln His Glu Leu Ser Pro Ser Thr Ser Thr Cys Lys
115 120 125

His Lys Val Gly Asp Thr Glu Gly Asp Tyr His Arg Ala Ser Ser Glu
130 135 140

Asn Cys Tyr Ser Val Tyr Ala Asp Gln Val Asn Ile Asp Tyr Leu Met
145 150 155 160

Asn Arg Pro Gln Asn Leu Arg Leu Glu Met Thr Ala Ala Lys Asn Thr
165 170 175

Asn Asn Asn Gln Ser Pro Ser Ala Pro Pro Ala Lys Pro Pro Ser Thr
180 185 190

Gln Arg Ala Val Ile Ser Pro Asp Gly Glu Cys Ser Ile Asp Asp Leu
195 200 205

Ser Phe Tyr Val Asn Arg Leu Ser Ser Leu Val Ile Gln Met Ala His
210 215 220

Lys Glu Ile Lys Glu Lys Leu Glu Gly Lys Ser Lys Cys Leu His His
225 230 235 240

Ser Ile Cys Pro Ser Pro Gly Asn Lys Glu Arg Ile Ser Pro Arg Thr
245 250 255

Pro Ala Ser Lys Ile Ala Ser Glu Met Ala Tyr Glu Ala Val Glu Leu
260 265 270

Thr Ala Ala Glu Met Arg Gly Thr Gly Glu Glu Ser Arg Glu Gly Gly
275 280 285

Gln Lys Ser Phe Leu Tyr Ser Glu Leu Ser Asn Lys Ser Lys Ser Gly
290 295 300

Asp Lys Gln Met Ser Gln Arg Glu Ser Lys Glu Phe Ala Asp Ser Ile
305 310 315 320

Ser Lys Gly Leu Met Val Tyr Ala Asn Gln Val Ala Ser Asp Met Met

09890709.012002

09890709.012662

Val Ser Leu	Met Lys Thr	Leu Lys	Val His Ser	Ser Gly Lys	Pro Ile
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Pro Ala Ser	Val Val Leu	Lys Arg	Val Leu Leu	Arg His Thr	Lys Glu
	355		360	365	
Ile Val Ser	Asp Leu Ile	Asp Ser Cys	Met Lys	Asn Leu His	Asn Ile
	370		375	380	
Thr Gly Val	Leu Met Thr	Asp Ser Asp	Phe Val Ser	Ala Val Lys	Arg
	385		390	395	400
Asn Leu Phe	Asn Gln Trp	Lys Gln Asn	Ala Thr Asp	Ile Met	Glu Ala
	405		410		415
Met Leu Ile	Leu Leu Val	Ser Ala Leu	Ile Gly Glu	Glu Lys	Glu Thr
	420		425	430	
Lys Ser Gln	Ser Leu Ser	Tyr Ala Ser	Leu Lys Ala	Gly Ser His	Asp
	435		440	445	
Pro Lys Cys	Arg Asn Gln	Ser Leu Glu	Phe Ser Thr	Met Lys Ala	Glu
	450		455	460	
Met Lys Glu	Arg Asp Lys	Gly Lys Met	Lys Ser Asp	Pro Cys Lys	Ser
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Leu Thr Ser	Ala Glu Lys	Val Gly Glu	His Ile Leu	Lys Glu Gly	Leu
	485		490	495	
Thr Ile Trp	Asn Gln Lys	Gln Gly Asn	Ser Cys Met	Val Ala Thr	Lys
	500		505	510	
Ala Cys Ser	Asn Lys Asp	Glu Lys Gly	Glu Lys Ile	Asn Ala Ser	Thr
	515		520	525	
Asp Ser Leu	Ala Lys Asp	Leu Ile Val	Ser Ala Leu	Lys Leu Ile	Gln
	530		535	540	
Tyr His Leu	Thr Gln Gln	Thr Lys Gly	Lys Asp Thr	Cys Glu Glu	Asp
	545		550	555	560
Cys Pro Gly	Ser Thr Met	Gly Tyr Met	Ala Gln Ser	Thr Gln Tyr	Glu
	565		570	575	
Lys Cys Gly	Gly Gly Gln	Ser Ala Lys	Ala Leu Ser	Val Lys Gln	Leu
	580		585	590	
Glu Ser His	Arg Ala Pro	Gly Pro Ser	Thr Cys Gln	Lys Glu Asn	Gln
	595		600	605	
His Leu Asp	Ser Gln Lys	Met Asp Met	Ser Asn Ile	Val Leu Met	Leu
	610		615	620	
Ile Gln Lys	Leu Leu Asn	Glu Asn Pro	Phe Lys Cys	Glu Asp Pro	Cys
	625		630	635	640
Glu Gly Glu	Asn Lys Cys	Ser Glu Pro	Arg Ala Ser	Lys Ala Ala	Ser
	645		650	655	
Met Ser Asn	Arg Ser Asp	Lys Ala Glu	Glu Gln Cys	Gln Glu His	Gln

660	665	670
Glu Leu Asp Cys Thr Ser Gly Met Lys Gln Ala Asn Gly Gln Phe Ile		
675	680	685
Asp Lys Leu Val Glu Ser Val Met Lys Leu Cys Leu Ile Met Ala Lys		
690	695	700
Tyr Ser Asn Asp Gly Ala Ala Leu Ala Glu Leu Glu Gln Ala Ala		
705	710	715
Ser Ala Asn Lys Pro Asn Phe Arg Gly Thr Arg Cys Ile His Ser Gly		
725	730	735
Ala Met Pro Gln Asn Tyr Gln Asp Ser Leu Gly His Glu Val Ile Val		
740	745	750
Asn Asn Gln Cys Ser Thr Asn Ser Leu Gln Lys Gln Leu Gln Ala Val		
755	760	765
Leu Gln Trp Ile Ala Ala Ser Gln Phe Asn Val Pro Met Leu Tyr Phe		
770	775	780
Met Gly Asp Lys Asp Gly Gln Leu Glu Lys Leu Pro Gln Val Ser Ala		
785	790	795
Lys Ala Ala Glu Lys Gly Tyr Ser Val Gly Gly Leu Leu Gln Glu Val		
805	810	815
Met Lys Phe Ala Lys Glu Arg Gln Pro Asp Glu Ala Val Gly Lys Val		
820	825	830
Ala Arg Lys Gln Leu Leu Asp Trp Leu Leu Ala Asn Leu		
835	840	845

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<211> 18

<212> PRT

<213> Homo sapiens

<400> 5

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1 5 10 15

Ile Asp

<210> 6

<211> 18

<212> PRT

<213> Homo sapiens

<400> 6

Gln Lys Ala Thr Asp Ile Met Asp Ala Met Leu Arg Lys Leu Tyr Asn
1 5 10 15

Val Met

<210> 7

<211> 18

<212> PRT

<213> Homo sapiens

<400> 7

Glu His Leu Met Asn Ser Val Met Lys Leu Cys Val Ile Ile Ala Lys
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Ser Cys

<210> 8

<211> 24

<212> DNA

<213> Artificial

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<223> primer

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<221> misc_feature

<223> N = inosine

<400> 8

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<213> Artificial

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